# Problems and Paradoxes in High-dimensional Data

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## Outline

- Motivation
- 2 Paradoxes
- Problems
- 4 A practical example
- Some positives
- Questions?

• Biotech: DNA microarrays, proteomics, transcriptomics etc.

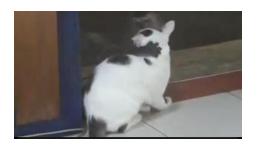
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## Examples

240p image  $\sim 100,000$  pixels, 3 color bandwidth  $p \approx 300,000$ 



• What's the volume of a circle with radius r?

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## Volume of *p*-dimensional ball with radius r > 0

$$V(p;r) = \frac{1}{\Gamma(p/2+1)} \pi^{p/2} r^p \tag{1}$$

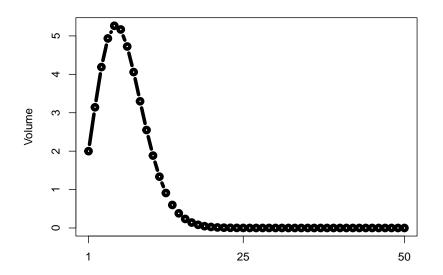
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Does the volume get bigger or smaller as we increase p?

Or more generally, how does the volume behave?

## The volume

## The volume



We've noticed that the volume of a *p*-dimensional ball tends to 0 as *p* increases, then naturally we might ask:

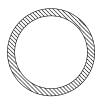
Where is the mass concentrated?

Consider the volume in the crust, i.e. the volume

$$C(p; r) = V(p; r) - V(p; 0.99r)$$
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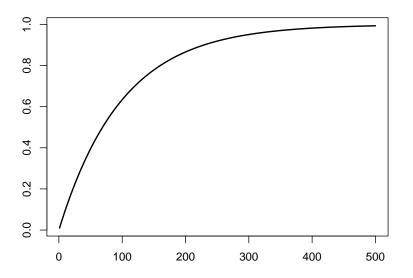
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Then the volume in the crust as a fraction of the total volume, i.e.

$$\frac{C(p;r)}{V(p;r)} = 1 - 0.99^p \tag{3}$$





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### Recap

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The lesson in all this is, we must be careful with our **geometric intuition** of high-dimensional spaces!

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- Most of the mass of a standard Gaussian is in the tail!
- Rare events may not actually be that rare

## **Problems**

# Data get far apart fast

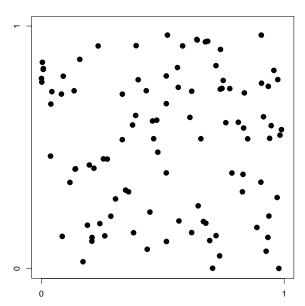
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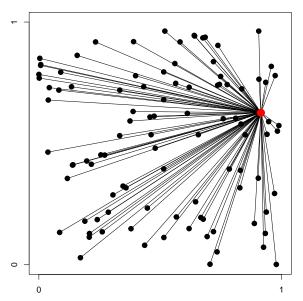
Suppose we have data uniformly distributed on a 2-D grid, 10-D grid, 100-D grid or 1000-D grid.

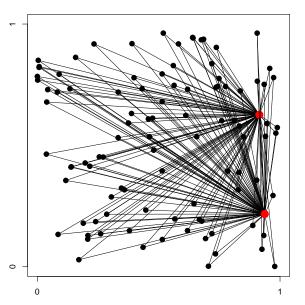
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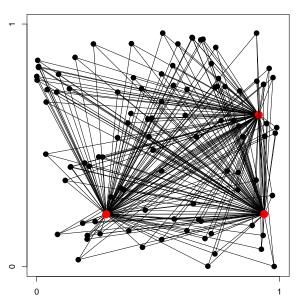
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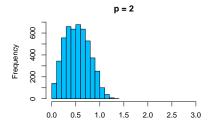
What's the average (euclidean) distance between points?

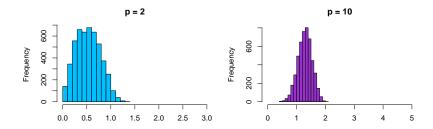


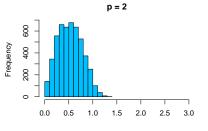


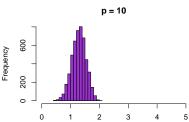


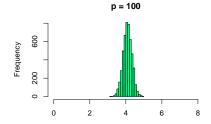


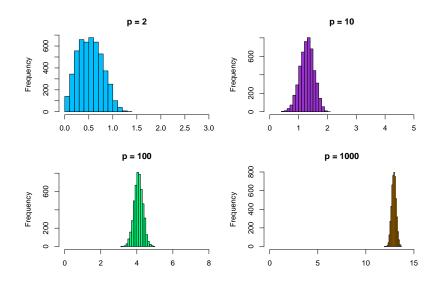


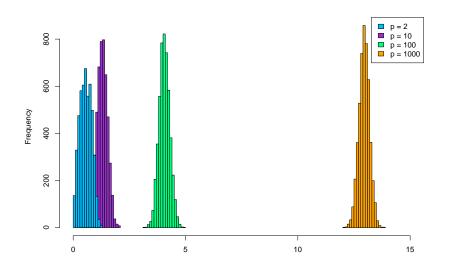












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- All points are a similar distance from the others
- The notion of nearest point vanishes

# Some more problems

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High-dimensional spaces are immense:

- Vast, data points can be isolated in the immensity
- Small changes add up fast. Many small fluctuations in different directions can produce a large global fluctuation
- Computation suffers

A practical example

PCA relies on the estimation of the covariance matrix  $\Sigma$ .

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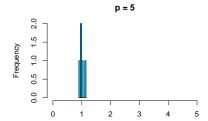
Let's suppose we have some some data generated iid from Normal(0,  $I_p$ ), where  $I_p$  is the  $p \times p$  identity.

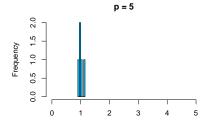
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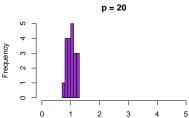
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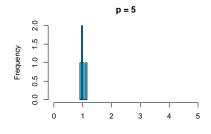
How well is  $\Sigma$  estimated as p grows?

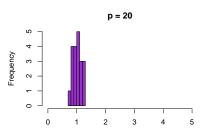
To see how well we're doing we're going to compare the eigenvalues of the estimated covariance matrices to that of the true covariance matrix  $(I_p)$ , we let n = 1000 and let p = 5, 20, 250, 500.

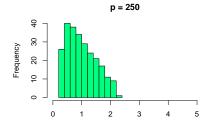


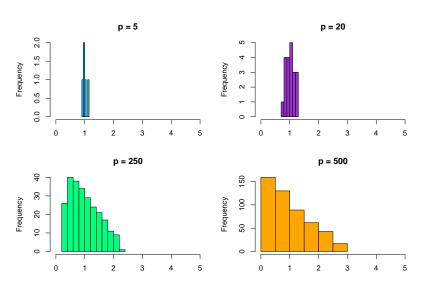












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In turn, downstream use of the covariance matrix (e.g. in PCA) can lead to spurious results.

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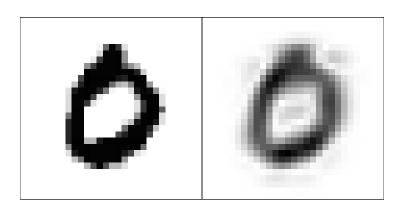
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### PCA on MNIST

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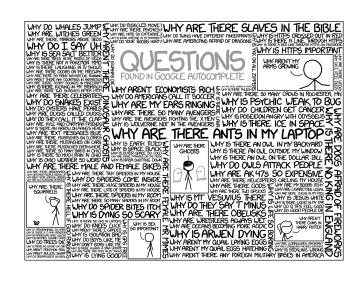
### Reconstructing the MNIST digits from the first 25 PCs



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Methods that can identify relevant structures or incorporate the knowledge that these spaces are often sparse generally outperform traditional methods.



#### Reference I

[Gir21] Christophe Giraud. *Introduction to High-Dimensional Statistics*. Chapman and Hall/CRC, Aug. 2021.